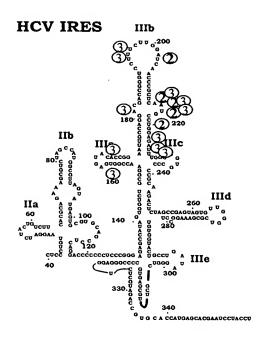
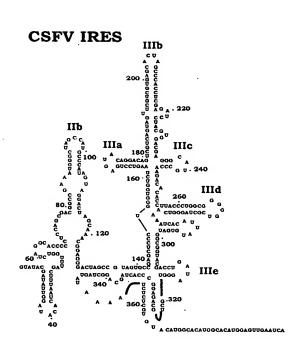
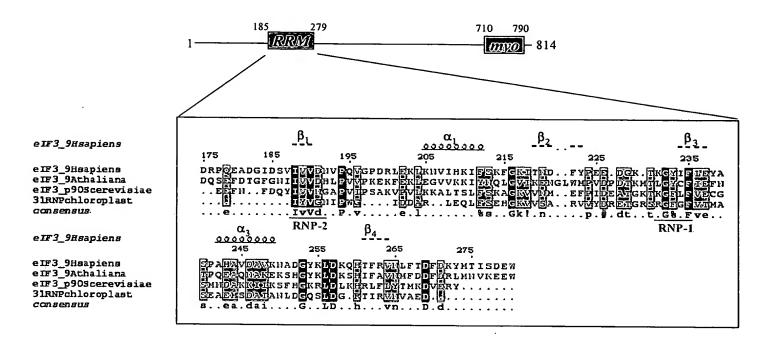
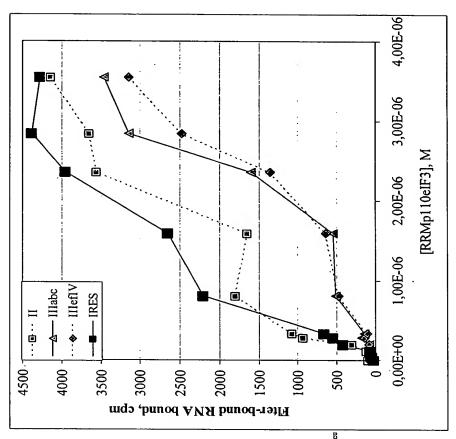
1/11

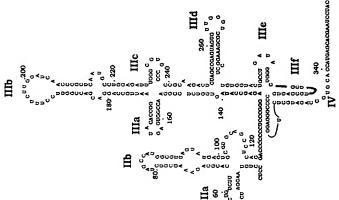












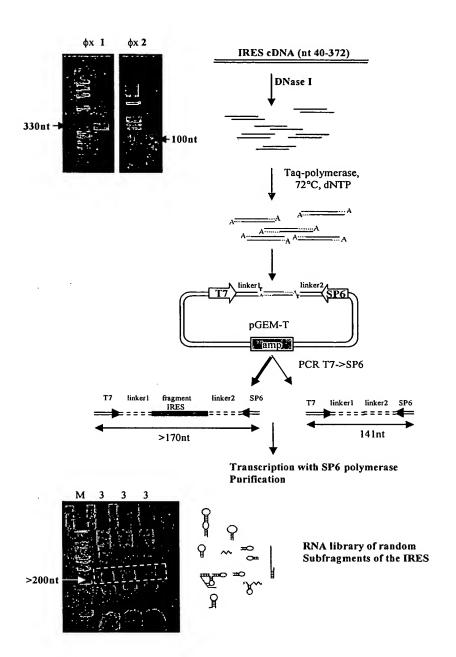


FIGURE 5A

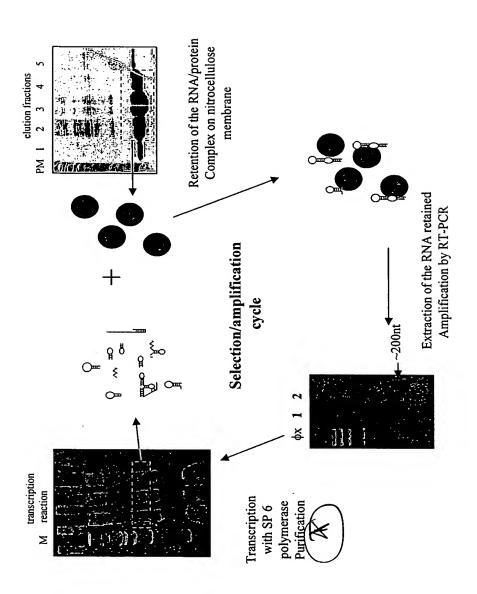


FIGURE 5B

Double- Strande	T7	linker l	fragment IRES	linker2	SP6					
DNA 3º	ATTATGCTGAGTGATATCCC	GCTTAAGCCCTTAANN	мининининининининининининининининининин	NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN	Beginning of transcription					
RNA 3	, v									
Single- Stranded, taatacgactcactatagggcgaattcgggaattnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnnn										
Double- Stranded DNA	₹TAATACGACTCACTATAGG ATTATGCTGAGTGATATCC	GCGAATTCGGGAATTNI CGCTTAAGCCCTTAANI	 	NNNNNNNNNNNNNNATCCAACGCGTTGGATGCATAG NNNNNNNNNNNNNTAGGTTGCGCAACCTACGTATC	Linker2-SP6 CTTGAGTATTCTATAGTGTCACCTAAATA3' GAACTCATAAGATATCACAGTGGATTTAT5'					

	10	20	30	40	50	60
	10	20	J U	70	- 20	90
1 10xx0	I	TACCCCTACCC	TCCCCT	 	ACTGT CTT	
4 28xx1				CTG AGGA ACT	ACTOT CITY	INC-CCA
4 35xx2					ACTGT-CTTC	
4 45xx3				ന്ന മനുവ	ACTOI-CITY	71C-GC 1
4 46xx4					ACTOI-CTTY	
5 2 xx x5			CCT	CTC ACCA ACT	ACTOC-CITY	CAC-GCA
5 4xxx6					ACTGT-CTTY	CICAGCA
5 6xxx7		TACCGCTAGCC	-тессет	CTG AGGA-CT	ACTGT-CTTY	LC-GCA
5 10xx8		TACCGCTGGCC	TCCCCT	CTG AGGA ACT	ACTGT-CTTY	CIC-GC A
5 12xx9					TACEAAGCGCC	LIGGOT
5 14xx10				T	LCTGT-CTTY	CIC-GCI
5 16x: 1.1						
HCV IRES	TTGGGGGCGACAC					
					*** *	t **
:						
	70	80	90	100	110	120
	l l	1	1	1	ı	1
4_10xx0	GAAAGCGT-CTAG					
4_28xx1	GAAAGCGT-CTAG					
4_3.5××2	GAA AGCGT-CTAG					
4_45xx3	GAAAGCGT-CTAG					
4_46xx4	GAAAGCGT-CTAG	CCATGGCGTT				
5_Z xx x5	GAA-GCGT-CTAG					
5 <u>4</u> xxx6	GAAAGCGT-CTAG					
5_6xxx7	GALAGOGIGOTAG					
5_10xx8	GAAAGCGT-CTAG	TTDDDDDTKDD	AGT L T	`CAG		
5_12xx9	AGACGCTTTCT	GCGTGAAGAC	agt l -			
5_14xx10	GAA-GCGT-CTAG					
5_16xx11	GAA-GCGT-CT-G					
ਸਨਾ IRES	GAA AGCGT-CTAG	CCATGGCGTT	ተፈጥብል ——-	<u>ጉ</u> ል ርብ የርብ ተ	FCAGCCȚCC A	GGACCCC

FIGURE 6A

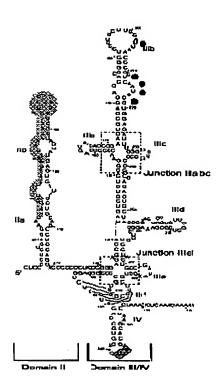
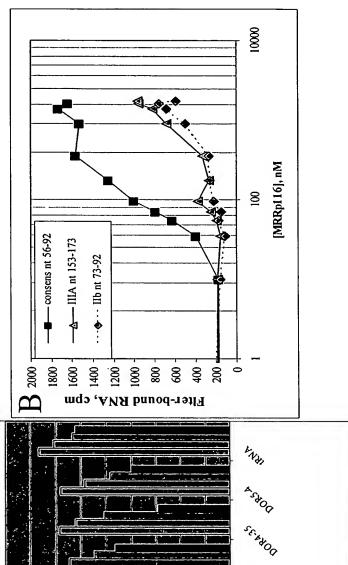


FIGURE 6B

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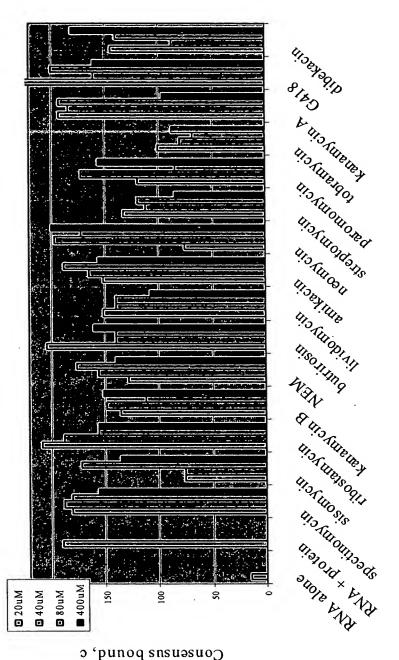
11

Mu2,0 ■

IRES bound to p110 RRM,cpm

Авоо в 0,2 µМ

Wisco New BROOK STAI



Consensus bound, c

